SEQ ID NO:9 alignment

```
<!--StartFragment-->RESULT 4
AAY79101
ΤD
    AAY79101 standard; protein; 108 AA.
XX
AC
    AAY79101;
XX
DT
    23-MAY-2000
                (first entry)
XX
DE
    Antibody light chain variable region.
XX
KW
    Antibody; light chain variable region; protein secretion; glycosylation.
XX
OS
    Unidentified.
XX
PΝ
    WO200005389-A2.
XX
PD
    03-FEB-2000.
XX
PF
    08-JUL-1999;
                  99WO-EP004919.
XX
PR
     20-JUL-1998;
                  98EP-00202432.
XX
PΑ
     (UNIL ) UNILEVER NV.
PA
     (UNIL ) UNILEVER PLC.
XX
PΙ
    Frenken LGJ, Sagt C, Verkleij AJ, Verrips CT;
XX
DR
    WPI; 2000-182710/16.
XX
PΤ
    High level recombinant production of heterologous protein with increased
PΤ
     secretory efficiency in lower eukaryotic cells, useful for producing
PT
    lysozyme, cutinase, or antibodies.
XX
PS
    Disclosure; Page 23; 55pp; English.
XX
CC
    The present sequence is that of the light chain variable region (VL) of
CC
    an antibody. The VL can be joined via a peptide linker to a heavy chain
CC
    variable region (VH) to form a single chain scFv fragment. The invention
CC
    relates to the high level recombinant production of proteins, such as
CC
    antibody fragments, in lower eukaryotic host (yeast or mould) cells. The
CC
    protein is N-qlycosylated at a location between its N-terminus and first
CC
    hydrophobic region to increase its level of secretion from the host cell
CC
    without affecting its functionality
XX
    Sequence 108 AA;
SO
 Query Match
                         100.0%; Score 550; DB 1; Length 108;
                         100.0%; Pred. No. 4.1e-33;
 Best Local Similarity
 Matches 103; Conservative
                             0; Mismatches
                                               0; Indels
                                                              0; Gaps
           1 ELTQSPASLSASVGETVTITCRASGNIHNYLAWYQQKQGKSPQLLVYYTTTLADGVPSRF 60
QУ
             3 ELTQSPASLSASVGETVTITCRASGNIHNYLAWYQQKQGKSPQLLVYYTTTLADGVPSRF 62
Db
          61 SGSGSGTQYSLKINSLQPEDFGSYYCQHFWSTPRTFGGGTKLE 103
QУ
             63 SGSGSGTQYSLKINSLQPEDFGSYYCQHFWSTPRTFGGGTKLE 105
<!--EndFragment-->
```